I,

```
\cdot_{\mathcal{Q}} Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Title:
.Perfect score:
sequence:
          Result
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OM nucleic - nucleic search, using sw model
                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
    Query
Score Match Length DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      September 3, 2002, 02:41:25; search time 2015.08 Seconds (without alignments) 228.469 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GenEmbl:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1797656 seqs, 10463268293 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 tgactgtgaacgttcgagatga 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-802-359B-1
22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gb_ba:*
gb_htg:*
gb_in:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gb_om:*
gb_ov:*
gb_pat:*
gb_ph:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gb_pr:*
                                                                                                                                         em_htg_hum:*
em_htg_inv:*
em_htg_other:*
em_htgo_inv:*
                                                                                                                                                                                                                                                                                                                                                                               gb_sts:*
gb_un:*
gb_vi:*
gb_vi:*
em_ba:*
em_fun:*
                                                                                                                                                                                                                                                                            em_pat:*
em_ph:*
                                                                                                                                                                                                                                                                                                        em_nu:*
em_om:*
em_or:*
                                                                                                                                                                                                                                               em_ro:*
                                                                                                                                                                                                                                                                                                                                                                   em_in:*
                                                                                                                                                                                                                     em_un:*
                                                                                                                                                                                                                                 em_sts:*
            ID
                                                       SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3595312
Description
```

	3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3		
0000000	222222222222222222222222222222222222	000	
92.7 92.7 92.7 92.7 92.7 91.8	922.77777777777777777777777777777777777		100.0 100.0 100.0 100.0 100.0 100.0 100.0
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0			
თთთთთთთ	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	<b>ຉຉຉຉຉຉຉຉຉ</b>	, <b>,</b> , , , , , , , , , , , , , , , , ,
5293 5311 5312 5312 5312 5252	AX083678 AX148639 AX148639 AX250702 AX250704 AX250708 AX252294 AX252294 AX252510 AX252512 AX252512 AX252512 AX252512 AX252523 AX25253	83694 8369 8369 83694 83694	3694 4699 3567 5070 5229 53111
293 311 311 312 312 312 252	XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX	25070 08368 114864 25229 25252 25252 25311 25311 26312 03694	0369 0469 1356 1356 1356 1350 150 150 150 150 150 150 150 150 150 1
Sequence Sequence Sequence Sequence Sequence Sequence	equencequencequencequencequencequencequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequenceequencee		Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence

## ALIGNMENTS

BASE COUNT ORIGIN	FEATURES source		AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	AX036945	RESULT 1
/organism="synthetic construct" /db_xref="taxon:32630" /note=" oligodesoxynucleotide" 6 a 3 c 7 g 6 t	Location/Qualifiers 122	ASSIST PUBL HOPITAUX DE PARIS (FR)	Carpentier, A.	1 (bases 1 to 22)	artificial sequence.	synthetic construct	synthetic construct.		AX036945.1 GI:11226373	AX036945	Sequence 2 from Patent FR2790955.	AX036945 22 bp DNA ]		
												linear		
												PAT 16-NOV-2000	×	(

ζ

```
VERSION
KEYWORDS
SOURCE
                                                                                                                                                                                                                                                                                      DEFINITION
ACCESSION
VERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACCESSION
                                                                                                                                                                                                                                                                                                                                    RESULT
AX083675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
AX046993
                                                                                      BASE COUNT
ORIGIN
   Ω<sub>γ</sub>
                                                                                                                                                                                                                                                                                                                                                                                                     õ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        δÃ
                                                                                                                                                          FEATURES
                                                                                                                                                                                                                              REFERENCE
                                                                                                                                                                                                                                                               SOURCE
                                                                                                                                                                                                                                                                                                                                                                                   밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE
                                                                                                                                                                                                                                                                            KEYWORDS
                                                                                                                                                                                                                                                                                                                         LOCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 22
                                                                                                                                                                                                      AUTHORS
TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AUTHORS
TITLE
                             Matches
                                     Query Match
Best Local
                                                                                                                                                                                 JOURNAL
                                                                                                                                                                                                                                                   ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM
                                                                                                                                                source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            source
                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                              1 tgactgtgaacgttcgagatga 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Н
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 tgactgtgaacgttcgagatga 22
tgactgtgaacgttcgagatga 22
                                                                                                                                                                                                                                                                                                                                                                                   TGACTGTGAACGTTCGAGATGA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGACTGTGAACGTTCGAGATGA 22
                                                                                                                                            van Nest,G.

Wethods of modulating an immune response using equences and compositions for use therein Patent: WO 011223-A 1 22-FEB-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                              22 bp
Sequence 1 from Patent W00112223.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AX046993 22 bp
Sequence 2 from Patent WO0067787.
AX046993
                                                                                                                                                                                                                                                  synthetic construct. synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hiv immunogenic compositions and methods Patent: WO 0067787-A 2 16-NOV-2000; THE IMMUNE RESPONSE CORPORATION (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  synthetic construct.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AX046993.1 GI:11876420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22;
                                                                                                                                                                                                                            artificial sequence.
1 (bases 1 to 22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Moss, R.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               artificial sequence.
1 (bases 1 to 22)
                                                                                                                                                                                                                                                                                       AX083675.1 GI:13185407
                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
llarity 100.0%;
Conservative (
                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="synthetic construct"
/db_xref="taxon:32630"
/note="phosphorothioate-modified
oligodeoxynucleotide"
a 3 c 7 g 6 t
                                                                                               /organism="synthetic construct"
/db_xref="taxon:33630"
/note="Synthetic construct"
3 c 7 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                               Score 22; DB
Pred. No. 0.2
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 22;
Pred. No.
                                       Score 22; E
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.25;
                                                                                                                                                                                                                                                                                                                                                                                                                                           0.25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       DΒ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
                                 <sub>уВ</sub> 6;
J.25;
                                                    DB
                                                                                                                                                                                                                                                                                                                          DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                      6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6;
                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                             0,
                                                  Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           synthetic
                             Indels
                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                       immunostimulatory s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PAT 15-DEC-2000
                                                                                                                                                                                                                                                                                                                          PAT 28-FEB-2001
                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                             Gaps
                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
```

```
REFERENCE
                                                                                                                                                                                                                                                                            VERSION
KEYWORDS
SOURCE
                                                                                                                                                                                                                                                                                                                                   RESULT
AX148636
LOCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KEYWORDS
SOURCE
ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 4
AX135650
LOCUS
DEFINITION
                                                                                                                                                                                                                                                                                                             DEFINITION
ACCESSION
  Вр
                                                                                                                                                                                                                                                                                                                                                                                             밁
                                                                                                                                                                                                                                                                                                                                                                                                                  Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
                      2
                                                                                                       BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE
                                                                                                                                                                         FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACCESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AUTHORS
TITLE
JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 22; Conservative
                                                        Query Match
Best Local
                                                                                                                                                                                                                     TITLE
                                                                                                                                                                                                                                                                    ORGANISM
                                                Matches
                                                                                                                                                                                               JOURNAL
                                                                                                                                                             source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         source
سر
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ...
                                                                                                                                                                                                                                                                                                                                                                                            tgactgtgaacgttcgagatga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGACTGTGAACGTTCGAGATGA
                                                                                                                                                                                                                             synthetic construct.
synthetic construct
artificial sequence.
1 (bases 1 to 22)
Tuck, S. and van Nest, G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AX135650
Sequence
AX135650
                                                                                                                                                                      Immunomodulatory compositions containing an immunostimulatory sequence linked to antigen and methods of use thereof Patent: WO 013591-A 1 25-MAY-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                             ch 100.0%; l Similarity 100.0%; 22; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cpg receptor (cpg-r) and methods relating thereto Patent: WO 0132877-A 21 10-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               artificial sequence.
1 (bases 1 to 22)
Mackichan, M.L.
                                                                                                                                                                                                                                                                                                    AX148636.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHIRON CORPORATION (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AX135650.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             σ
                                                                                                                  σ
                                                                                                                  Ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                 /organism="synthetic construct"
/db_xref="taxon:32630"
/note="synthetic construct"
/note="synthetic construct"
3 c 7 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="CpG oligonucleotide" 3 c 7 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="synthetic construct"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22 bp
from Patent W00132877.
                                                                                                                                                                                                                                                                                                     GI:14347254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GI:14271920
                                                0;
                                                                                                                                                                                                                                                                                                                                                                                              22
                                                                                                                                                                                                                                                                                                                                                                                                                   22
                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22
                                              Score 22; DB 6
Pred. No. 0.25;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 22;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 6; Length 22; 0.25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA
                                                                                                                                                                                                                                                                                                                                      DNA
                                                                       6;
                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                    Length 22;
                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PAT 29-MAY-2001
                                                                                                                                                                                                                                                                                                                                       PAT 08-6UN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                 Gaps
                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                            0
```

RESULT AX250701

```
ċ
                                                                                                                                                                                                                                                                                                                                                                                   DEFINITION
ACCESSION
VERSION
KEYWORDS
VERSION
KEYWORDS
SOURCE
                                 DEFINITION ACCESSION
                                                               RESULT
AX252509
                                                                                                           В
                                                                                                                         δõ
                                                                                                                                                                                                         BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT
AX252291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE
AUTHORS
TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCUS
DEFINITION
ACCESSION
                                                                                                                                                                                                                                                                                                                                          REFERENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORIGIN
                                                      LOCUS
                                                                                                                                                                                                                                                                          FEATURES
                                                                                                                                                                                                                                                                                                                                                                          SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BASE COUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                              rocus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOURCE
ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KEYWORDS
                                                                                                                                                    Query Match
Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VERSION
                                                                                                                                                                                                                                                                                                                    AUTHORS
TITLE
                                                                                                                                                                                                                                                                                                                                                               ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JOURNAL
                                                                                                                                                                                                                                                               source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            source
                                                                                                           _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 tgactgtgaacgttcgagatga 22
                                                                            œ
                                                                                                                     tgactgtgaacgttcgagatga 22
                                                                                                       TGACTGTGAACGTTCGAGATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TGACTGTGAACGTTCGAGATGA
                                                                                                                                                                                                                                                                                                                                                                                                      22 bp
Sequence 1 from Patent WO0168117.
AX252291
                                 AX252509
Sequence 1 from
AX252509
                                                                                                                                                     . Similarity
22; Conserv
                                                                                                                                                                                                                                                                               polynucleotide sequences
Patent: WO 0168117-A 1 20-SEP-2001;
Dynavax Technologies Corporation (U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
22; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent:
Dynavax
 synthetic
                      AX252509.1
                                                                                                                                                                                                                                                                                                                                                             synthetic construct
                                                                                                                                                                                                                                                                                                                                                                          synthetic construct.
                                                                                                                                                                                                                                                                                                                Methods of reducing papillomavirus infection using immunomodulatory
                                                                                                                                                                                                                                                                                                                               van Nest, G.
                                                                                                                                                                                                                                                                                                                                                                                             AX252291.1 GI:15985632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          synthetic construct synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AX250701 22 bp
Sequence 1 from Patent WO0168078.
AX250701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          van Nest,G.
Methods of suppressing hepatitis virus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AX250701.1 GI:15984439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      artificial sequence.
1 (bases 1 to 22)
                                                                                                                                                                                                                                                                                                                                        (bases 1 to 22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
llarity 100.0%;
Conservative 0;
                                                                                                                                                     Conservative
                                                                                                                                                                                                                   σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6
                                                                                                                                                                                                                 /organism="synthetic construct"
/dh_xref="taxon:32630"
/note="Polynucleotide containing CG"
7 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="synthetic construct"
/db_xref="taxon:32630"
3 c 7 g 6 t
                                                                                                                                                                                                                                                                        Location/Qualifiers
 construct
                      GI:15985780
                                                                                                                                                             100.0%;
                                       22 bp
Patent WO0168103
                                                                                                           22
                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 22; DE
Pred. No. 0.2
); Mismatches
                                                                                                                                                   Score 22; DB 6;
Pred. No. 0.25;
; Mismatches (
                                                                                                                                                                                                                                                                                 (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 6;
0.25;
                                                                                                                                                                                                                                                                                                                                                                                                                             DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA
                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 22;
                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 using
                                                    PAT 05-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                            PAT 05-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PAT 06-OCT-2001
                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
```

```
RESULT 1
AX252934
                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE
AUTHORS
TITLE
                                                                     VERSION
KEYWORDS
                                                                                                      LOCUS
DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KEYWORDS
SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DEFINITION
ACCESSION
VERSION
                                                                                                                                                                              밁
                                                                                                                                                                                                 ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT
AX252520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE
AUTHORS
TITLE
                       REFERENCE
                                                           SOURCE
                                                                                                                                                                                                                                                                                     ORIGIN
                                                                                                                                                                                                                                                                                               BASE COUNT
                                                                                                                                                                                                                                                                                                                                                          FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURES
                                                                                            ACCESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BASE COUNT
                                                                                                                                                                                                                        Query Match
Best Local S
Matches 22
                                               ORGANISM
 TITLE
            AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                 JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM
                                                                                                                                                                                                                                                                                                                                               source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      source
                                                                                                                                         10
                                                                                                                                                                              μ.
                                                                                                                                                                                       tgactgtgaacgttcgagatga 22
                                                                                                                                                                            TGACTGTGAACGTTCGAGATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGACTGTGAACGTTCGAGATGA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tgactgtgaacgttcgagatga 22
                     artificial sequence.
1 (bases 1 to 22)
                                             synthetic construct
                                                                                           Sequence 1
AX252934
                                                                                                                                                                                                                         22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                               artificial sequence.
1 (bases 1 to 22)
van Nest,G. and Tuck,S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               van Nest, G. and Tuck, S.
                                                       synthetic construct
                                                                                AX252934.1
                                                                                                                 AX252934
                                                                                                                                                                                                                                                                                                                                                       Patent: WO 0168144-A 1 20-SEP-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          synthetic construct
artificial sequence.
1 (bases 1 to 22)
van Nest.G.
Methods of ameliorating symptoms of herpes
                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AX252520.1
Immunomodulatory formulations
                                                                                                                                                                                                                                                                                                                                                                                                       Biodegradable immunomodulatory formulations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 synthetic construct.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunoodulatory polynucleotide sequences Patent: WO 0168103-A 1 20-SEP-2001; Dynavax Technologies Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                                              thereo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                       100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; illarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                δ
                                                                                                                                                                                                                                                                                                Ø
                                                                                                                                                                                                                                                                                           /organism="synthetic construct"
/db_xref="taxon:32630"
/note="Polynucleotide containing
3 c 7 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="synthetic construct"
/db_xref="taxon:32630"
/note="polynucleoctide containing
/note="polynucleoctide containing
3 c 7 g 6 t
                                                                                                     22 bp
from Patent W00168143
                                                                                GI:15986201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GI:15985791
                                                                                                                                                                            22
                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                        Score 22; DB 6;
Pred. No. 0.25;
; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NO.
and methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA
                                                                                                                 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6;
                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 22;
for
                                                                                                                                                                                                                                                                                                           င္ပ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  င္ပ
                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infection using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                      and methods for use
use thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PAT
                                                                                                                 PAT 05-OCT-2001
                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-OCT -2001
                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
```

```
ξ.
                                                                             REFERENCE
AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE
AUTHORS
TITLE
                                                                                                                                                  DEFINITION
ACCESSION
VERSION
                                                                                                                                                                                                                                                                                                                                                   BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BASE COUNT
ORIGIN
                                                                                                                           KEYWORDS
SOURCE
                                                                                                                                                                                               RESULT 12
AX253123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOURCE
ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1
AX253113
                                                                                                                                                                                     LOCUS
                                                                                                                                                                                                                                            밁
                                                                                                                                                                                                                                                              QΥ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             δõ
                      FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  POCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KEYWORDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACCESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 22
                                                                    TITLE
                                                                                                                 ORGANISM
                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                         Matches
                                              JOURNAL
             source
                                                                                                                                                                                                                                                                                                                                                                                                           source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 tgactgtgaacgttcgagatga 22
                                                                                                                                                                                                                                                         tgactgtgaacgttcgagatga 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGACTGTGAACGTTCGAGATGA 22
                                                                                                                                                                                                                                             TGACTGTGAACGTTCGAGATGA 22
                                                                            synthetic construct artificial sequence.
1 (bases 1 to 22)
van Nest, G.
                     Methods of preventing and treating viral immunodulatory polynucleotide sequences Patent: WO 0168077-A 1 20-SEP-2001; Dynavax Technologies Corporation (US) Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                   Methods of preventing and treating respiratory viral infection using immunomodulatory polynucleotide sequences
Patent: WO 0168116-A 1 20-SEP-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
                                                                                                                                                              AX253123 22 bp
Sequence 1 from Patent WO0168077.
AX253123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1
AX253113
                                                                                                                            synthetic construct.
                                                                                                                                                    AX253123.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AX253113.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AX253113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dynavax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            van Nest,G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                artificial sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (bases 1 to 22)
                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO 0168143-A 1 20-SEP-2001; Technologies Corporation (U
                                                                                                                                                                                                                                                                                                                                                          /organism="synthetic construct"
/db_xref="taxon:32630"
/note="polynucleotide containing
/ g c 7 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism≈"synthetic construct"
/db_xref∈"taxon:32630"
organism="synthetic construct"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     construct.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Polynucleotide containing
3 c 7 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     from Patent WO0168116.
                                                                                                                                                    GI:15986291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GI:15986281
                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 22;
Pred. No.
                                                                                                                                                                                                                                                                                                    Score 22; DB 6
Pred. No. 0.25;
                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22 bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DΒ
                                                                                                                                                                                                                                                                                                               6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6.
                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 22;
                                                                     infections
                                                                                                                                                                                                                                                                                                              Length 22;
                                                                                                                                                                                                                                                                                                                                                                        G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G,
                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                     PAT
                                                                    using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PAT
                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                     05-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KEYWORDS
SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 13
BD009235
LOCUS
                                                                    KEYWORDS
SOURCE
                                                                                                     DEFINITION ACCESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACCESSION
VERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BASE COUNT
ORIGIN
                                                                                                                                                                                     ₽
                                                                                                                                                                                                         Ş
                                                                                                                                                                                                                                                                                         BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Š
                                     REFERENCE
                                                                                                                            LOCUS
                                                                                                                                                                                                                                                                                                                                                   FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMMENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DEFINITION
                                                                                                                                      AX250707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE
                                                                                            VERSION
                                                                                                                                                   RESULT
                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 22
             AUTHORS
TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AUTHORS
TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local :
                                                         ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM
                                                                                                                                                                                                                                                                                                                                       source
                       van
```

```
1 tgactgtgaacgttcgagatga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         artificial sequence.
El (bases 1 to 22)
El (bases 1 to 22)
Immunostimulatory polynucleotide/immunomodulatory molecule
Immunostimulatory polynucleotide/immunomodulatory molecule
Patent: JP 2001503254-A 34 13-MAR-2001;
THE REGENTS OF THE UNIVERSITY OF CALIFORNIA
OS Artificial Sequence
PN 19-001503254-A/34
PD 13-MAR-2001
PP 09-0CT-1997 JP 1998518649
PF 10-0CT-1996 US 60/028118
PI 1-0CT-1996 US 60/028118
PI DENNIS A CARSON, EYAL RAZ, MARK ROMAN
PC A61K39/00, A61K39/385, A61K39/39
CC
FH Key
FT Source
/Organism='Artificial Sequence'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 TGACTGTGAACGTTCGAGATGA
                                              synthetic construct
synthetic construct
artificial sequence.
1 (bases 1 to 22)
                                                                                                                                     22 bp
Sequence 7 from Patent W00168078.
AX250707
                                                                                                                                                                                                                                                                                                          ch 100.0%;
l Similarity 100.0%;
22; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic construct. synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BD009235 22 bp DNA linear Immunostimulatory polynucleotide/immunomodulatory
                                                                                                                        AX250707.1
Methods of suppressing hepatitis virus infection using immunomodulatory polynucleotide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP 2001503254-A/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BD009235.1 GI:18637608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BD009235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   conjugates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                               Nest, G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6
                                                                                                                                                                                                                                                                                                                                                                                                    σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Polynucleotide a 3 c 7 g
                                                                                                                                                                                                                                                                                                                                                                                                    þ
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism='Artificial
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                   /organism="synthetic co
/db_xref="taxon:32630"
3 c 7 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:32630"
                                                                                                                          GI:15984445
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22
                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                          Score 22; DB
Pred. No. 0.2
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 22; Depred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 construct"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             containing
6 t
                                                                                                                                                                                                                                                                                                                       DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                    σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . 25;
                                                                                                                                                                      DNA
                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                      Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PAT 31-JAN-2002
                                                                                                                                                                      PAT 05-OCT-2001
                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                           0.
```

```
SOURCE
ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 15
AX083681
LOCUS
DEFINITION
ACCESSION
VERSION
                                                                                                                                                                         BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                                                                      REFERENCE
AUTHORS
TITLE
Search completed: September 3, 2002, 04:24:03 Job time: 6158 sec
                                                             В
                                                                                Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                           FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                 KEYWORDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURES
                                                                                                         Query Match 95.5%;
Best Local Similarity 95.5%;
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 96.4
Best Local Similarity 95.5
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                  JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JOURNAL
                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                            source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       source
                                                           synthetic construct.
synthetic construct
artificial sequence.
1 (bases 1 to 22)
van Nest,G.
Methods of modulating an immune response using immunostimulatory s
equences and compositions for use therein
Patent: WO 011223-A 7 22-FEB-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                  22 bp
Sequence 7 from Patent W00112223.
AX083681
AX083681.1 GI:13185413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent: WO 0168078-A 7 20-SEP-2001;
Dynavax Technologies Corporation (US)
Location/Qualifiers
                                                                                                                                                                                      6
a
                                                                                                                                                                                                                       /organism="synthetic construct"
/db_xref="taxon:32630"
11
                                                                                                                                                                                     /note="5-bromocytosine"
/mod_base=OTHER
1 2 c 7 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="synthetic construct"
/db_xxef="taxon:32630"
/note="B is 5-bromocytosine"
2 c 7 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96.4%;
95.5%;
                                                                                                                      Score 21; DB 6; Length 22; Pred. No. 0.9;
                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 21.2; DB 6; Length 22; Pred. No. 0.69;
                                                                                                                                                                                  7 9
                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                      6
t
                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                      1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 others
                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                PAT 28-FEB-2001
                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
```

THIS PAGE BLANK (USPTO)